

0280

OIKE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/008,093

DATE: 12/18/2001

TIME: 16:07:52

Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\12182001\J008093.raw

ENTERED

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3 <110> APPLICANT: McCarthy, Jeannette Just
5 <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE IDENTIFICATION,
6 ASSESSMENT, PREVENTION AND THERAPY OF
7 CARDIOVASCULAR DISEASE
9 <130> FILE REFERENCE: MRI-026
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/008,093
C--> 11 <141> CURRENT FILING DATE: 2001-11-09
11 <150> PRIOR APPLICATION NUMBER: US 60/248,185
12 <151> PRIOR FILING DATE: 2000-11-09
14 <150> PRIOR APPLICATION NUMBER: US 60/257,417
15 <151> PRIOR FILING DATE: 2000-12-22
17 <160> NUMBER OF SEQ ID NOS: 6
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22 <211> LENGTH: 5724
23 <212> TYPE: DNA
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58 atccagtgca cagatgttga tgagtgc aaa gaagtgcctg atgcctgctt caaccacaat 1920
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103 cttttttcct gaattatcat ggagttttct aattctctct tttggaatgt agattttttt 4620
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124 <210> SEQ ID NO: 2

125 <211> LENGTH: 1170

126 <212> TYPE: PRT

127 <213> ORGANISM: Homo sapiens

129 <400> SEQUENCE: 2

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133 20 25 30
134 Ile Phe Glu Leu Thr Gly Ala Ala Arg Lys Gly Ser Gly Arg Arg Leu
135 35 40 45
136 Val Lys Gly Pro Asp Pro Ser Ser Pro Ala Phe Arg Ile Glu Asp Ala
137 50 55 60
138 Asn Leu Ile Pro Pro Val Pro Asp Asp Lys Phe Gln Asp Leu Val Asp
139 65 70 75 80
140 Ala Val Arg Ala Glu Lys Gly Phe Leu Leu Ala Ser Leu Arg Gln
141 85 90 95
142 Met Lys Lys Thr Arg Gly Thr Leu Leu Ala Leu Glu Arg Lys Asp His
143 100 105 110
144 Ser Gly Gln Val Phe Ser Val Val Ser Asn Gly Lys Ala Gly Thr Leu
145 115 120 125
146 Asp Leu Ser Leu Thr Val Gln Gly Lys Gln His Val Val Ser Val Glu
147 130 135 140
148 Glu Ala Leu Leu Ala Thr Gly Gln Trp Lys Ser Ile Thr Leu Phe Val
149 145 150 155 160
150 Gln Glu Asp Arg Ala Gln Leu Tyr Ile Asp Cys Glu Lys Met Glu Asn
151 165 170 175
152 Ala Glu Leu Asp Val Pro Ile Gln Ser Val Phe Thr Arg Asp Leu Ala
153 180 185 190
154 Ser Ile Ala Arg Leu Arg Ile Ala Lys Gly Gly Val Asn Asp Asn Phe
155 195 200 205

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159 225      230      235      240
160 Thr Leu Asp Asn Asn Val Val Asn Gly Ser Ser Pro Ala Ile Arg Thr
161      245      250      255
162 Asn Tyr Ile Gly His Lys Thr Lys Asp Leu Gln Ala Ile Cys Gly Ile
163      260      265      270
164 Ser Cys Asp Glu Leu Ser Ser Met Val Leu Glu Leu Arg Gly Leu Arg
165      275      280      285
166 Thr Ile Val Thr Thr Leu Gln Asp Ser Ile Arg Lys Val Thr Glu Glu
167      290      295      300
168 Asn Lys Glu Leu Ala Asn Glu Leu Arg Arg Pro Pro Leu Cys Tyr His
169 305      310      315      320
170 Asn Gly Val Gln Tyr Arg Asn Asn Glu Glu Trp Thr Val Asp Ser Cys
171      325      330      335
172 Thr Glu Cys His Cys Gln Asn Ser Val Thr Ile Cys Lys Lys Val Ser
173      340      345      350
174 Cys Pro Ile Met Pro Cys Ser Asn Ala Thr Val Pro Asp Gly Glu Cys
175      355      360      365
176 Cys Pro Arg Cys Trp Pro Ser Asp Ser Ala Asp Asp Gly Trp Ser Pro
177      370      375      380
178 Trp Ser Glu Trp Thr Ser Cys Ser Thr Ser Cys Gly Asn Gly Ile Gln
179 385      390      395      400
180 Gln Arg Gly Arg Ser Cys Asp Ser Leu Asn Asn Arg Cys Glu Gly Ser
181      405      410      415
182 Ser Val Gln Thr Arg Thr Cys His Ile Gln Glu Cys Asp Lys Arg Phe
183      420      425      430
184 Lys Gln Asp Gly Gly Trp Ser His Trp Ser Pro Trp Ser Ser Cys Ser
185      435      440      445
186 Val Thr Cys Gly Asp Gly Val Ile Thr Arg Ile Arg Leu Cys Asn Ser
187      450      455      460
188 Pro Ser Pro Gln Met Asn Gly Lys Pro Cys Glu Gly Glu Ala Arg Glu
189 465      470      475      480
190 Thr Lys Ala Cys Lys Lys Asp Ala Cys Pro Ile Asn Gly Gly Trp Gly
191      485      490      495
192 Pro Trp Ser Pro Trp Asp Ile Cys Ser Val Thr Cys Gly Gly Gly Val
193      500      505      510
194 Gln Lys Arg Ser Arg Leu Cys Asn Asn Pro Ala Pro Gln Phe Gly Gly
195      515      520      525
196 Lys Asp Cys Val Gly Asp Val Thr Glu Asn Gln Ile Cys Asn Lys Gln
197      530      535      540
198 Asp Cys Pro Ile Asp Gly Cys Leu Ser Asn Pro Cys Phe Ala Gly Val
199 545      550      555      560
200 Lys Cys Thr Ser Tyr Pro Asp Gly Ser Trp Lys Cys Gly Ala Cys Pro
201      565      570      575
202 Pro Gly Tyr Ser Gly Asn Gly Ile Gln Cys Thr Asp Val Asp Glu Cys
203      580      585      590
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209 625          630          635          640
210 Lys Gln Val Cys Lys Pro Arg Asn Pro Cys Thr Asp Gly Thr His Asp
211          645          650          655
212 Cys Asn Lys Asn Ala Lys Cys Asn Tyr Leu Gly His Tyr Ser Asp Pro
213          660          665          670
214 Met Tyr Arg Cys Glu Cys Lys Pro Gly Tyr Ala Gly Asn Gly Ile Ile
215          675          680          685
216 Cys Gly Glu Asp Thr Asp Leu Asp Gly Trp Pro Asn Glu Asn Leu Val
217          690          695          700
218 Cys Val Ala Asn Ala Thr Tyr His Cys Lys Lys Asp Asn Cys Pro Asn
219 705          710          715          720
220 Leu Pro Asn Ser Gly Gln Glu Asp Tyr Asp Lys Asp Gly Ile Gly Asp
221          725          730          735
222 Ala Cys Asp Asp Asp Asp Asn Asp Lys Ile Pro Asp Asp Arg Asp
223          740          745          750
224 Asn Cys Pro Phe His Tyr Asn Pro Ala Gln Tyr Asp Tyr Asp Arg Asp
225          755          760          765
226 Asp Val Gly Asp Arg Cys Asp Asn Cys Pro Tyr Asn His Asn Pro Asp
227          770          775          780
228 Gln Ala Asp Thr Asp Asn Asn Gly Glu Gly Asp Ala Cys Ala Ala Asp
229 785          790          795          800
230 Ile Asp Gly Asp Gly Ile Leu Asn Glu Arg Asp Asn Cys Gln Tyr Val
231          805          810          815
232 Tyr Asn Val Asp Gln Arg Asp Thr Asp Met Asp Gly Val Gly Asp Gln
233          820          825          830
234 Cys Asp Asn Cys Pro Leu Glu His Asn Pro Asp Gln Leu Asp Ser Asp
235          835          840          845
236 Ser Asp Arg Ile Gly Asp Thr Cys Asp Asn Asn Gln Asp Ile Asp Glu
237          850          855          860
238 Asp Gly His Gln Asn Asn Leu Asp Asn Cys Pro Tyr Val Pro Asn Ala
239 865          870          875          880
240 Asn Gln Ala Asp His Asp Lys Asp Gly Lys Gly Asp Ala Cys Asp His
241          885          890          895
242 Asp Asp Asp Asn Asp Gly Ile Pro Asp Asp Lys Asp Asn Cys Arg Leu
243          900          905          910
244 Val Pro Asn Pro Asp Gln Lys Asp Ser Asp Gly Asp Gly Arg Gly Asp
245          915          920          925
246 Ala Cys Lys Asp Asp Phe Asp His Asp Ser Val Pro Asp Ile Asp Asp
247          930          935          940
248 Ile Cys Pro Glu Asn Val Asp Ile Ser Glu Thr Asp Phe Arg Arg Phe
249 945          950          955          960
250 Gln Met Ile Pro Leu Asp Pro Lys Gly Thr Ser Gln Asn Asp Pro Asn
251          965          970          975
252 Trp Val Val Arg His Gln Gly Lys Glu Leu Val Gln Thr Val Asn Cys
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VERIFICATION SUMMARY

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L:11 M:270 C: Current Application Number differs, Replaced Current Application No

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date